

# Multi-local evaluation of the agronomic performances of new sorghum lines in the villages of Riadi, Garin Maigari and on station at CERRA-Maradi

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## ABSTRACT

The field trial was conducted in 2021 rainy season at RIADI (13°25'47" north latitude and 7°7'33" east longitude, GARIN MAIGARI (13°36'88" latitude North and 7°08'86" East longitude) and the CERRA-Maradi STATION (13°27'72" North latitude and 7°06'54" East longitude) in order to evaluate the agronomic performance of new sorghum genotypes and to identify among them the best adapted genotypes in term of grain yields. The experiment contains thirty (30) different varieties including 24 varieties resulting from the cross between MDK and L153-5 genotypes, two parents (MDK and L-153-5), two station checks (L28 and SSD 35) and two local checks in each location. The treatments were laid out in an incomplete randomized block design with three replicates in each location. The results revealed positive and negative correlations between some variables such as RDT G and RDT T, RDT G and PP, RDT G and NPR, RDT G and HTR. The combined ANOVA revealed significant effects of genotype (G), environment (E) and genotype x environment interactions (GxE). In addition, GGE Biplot analysis, highly performed genotypes in grain yield such as V68, V3 and V58 were identified among the genotypes. Some genotypes have a specific adaptation, these are the MDK, TL2G, L28 and V5 genotypes which are the best genotypes in the mega environment formed by GARIN MAIGARI and RIADI and V27 and V73 which are the best genotypes in RIADI. This analysis allowed also to identify genotypes such as SSD 35, V61, L28, V27, MDK which have medium unstable yield means and V13, V 47, V 43, V19, V29, V8 which are stable but produced lower yields (below average). It also showed a correlation between the STATION and GARIN MAIGARI which constitute a mega environment and showed that GARIN MAIGARI is the ideal environment (reference). So, it can be used for genotypes with wide adaptation while non-representative environments such as RIADI and STATION are suitable for selecting specifically adapted genotypes.

**Keywords:** sorghum, multi-local evaluation, new genotypes, agronomic performances.

## I. INTRODUCTION

A landlocked Sahelian country with a growing population, Niger is one of the driest countries in the world where it rains around three to four months a year. This country is experiencing difficulties in ensuring its food self-sufficiency.

Agriculture and livestock farming alone employ 85% of the active population and contribute to the national economy with a GDP of 43% (INS, 2014a). With an area of 1,267,000 km<sup>2</sup>, Niger where agriculture occupies an important place in the socio-economic life of the populations, Niger has a large area of land of which only the southern strip (12%) constitutes a fertile and benefits from sufficient annual rainfall for the various crops without irrigation. Millet and sorghum are the main cereals cultivated both in terms of area sown, production and their contribution to food. These two cereals are cultivated in all regions of the country and occupy a privileged place except in Aïr (Idrissa, 2001, cited by Sadi, 2011). The place of sorghum is reflected above all by a considerable increase in sown areas which went from 2,544,700 ha in 2009 to more than 3,565,900 ha in 2013 (INS, 2014b). National production for the same period increased from 738,500 tons to 1,320,000 tons. Unfortunately, sorghum productivity remains very low, the national yield is equivalent to 343 kg/ha (INS, 2014b).

Agricultural production in Niger, as in the Sahel countries in general, is dependent on several factors, including the low fertility of cultivated land and the often deficit and irregular rainfall. In addition to these factors, sorghum production is limited by seed quality. The latter are obtained from traditional varieties generally with a long cycle. The grains matured during a period of high humidity. This hygrometry favors the establishment of pathogens (mold) altering the quality of sorghum seeds (Louvel, 1982 and 1983, cited by Louis, 1995).

In order to remedy this situation, several studies have been carried out, particularly in the context of the fight against drought which frequently occurs towards the end of crop cycles. Sorghum varieties better adapted to drought conditions have also been developed and popularized by research institutions such as INRAN and ICRISAT.

Despite these efforts, the main aim of which is to complement local varieties sensitive to the climatic conditions of the Sahel and to crop enemies with improved varieties adapted to these conditions, the use of local varieties remains the most dominant. This is explained by the unavailability of improved seeds among producers and the problem of choice of variety (Bassirou, 2014).

As a result, there is a growing need to increase sorghum productivity in Niger in response to these challenges and yield declines. To overcome this problem, a population resulting from the cross between MDK and L153-5 was developed in order to provide farmers with productive varieties adapted to the drastic conditions of Niger where among actions in favor of food security include the creation and/or introduction of new sorghum genotypes adapted to various environments in terms of soil and climate. It is in this same vein that this present work is carried out. The main objective of this work is a multi-local evaluation of the agronomic performances of new sorghum lines resulting from the crossing between MDK and L153-5 with the aim of identifying the best adapted lines with good productivity.

## II. MATERIALS AND METHODS

### 1. Plant material

This study includes thirty (30) lines of sorghum including 24 lines resulting from the cross between Matché Da Koumgna (MDK) a very productive late variety (3 t/ha) and L 153-5 a very early variety with low yield, two (2) lines from the INRAN sorghum program (L 28 and SSD 35) and two control varieties widely used locally at each of the three evaluation sites. Mota Maradi (MM) and El Rourouka for the INRAN Maradi research station site, Makaho da Wayo (MDW) and the Hakorin Karouwa (HK) variety for the Riadi site and finally the El Bazanga and MDW for the Garin Maigari site were used in this experiment.

### 2. Experimental site

The various experiments were carried out at three (3) different sites, namely the villages of Garin Maigari and Riadi and in the CERRA Maradi research station.

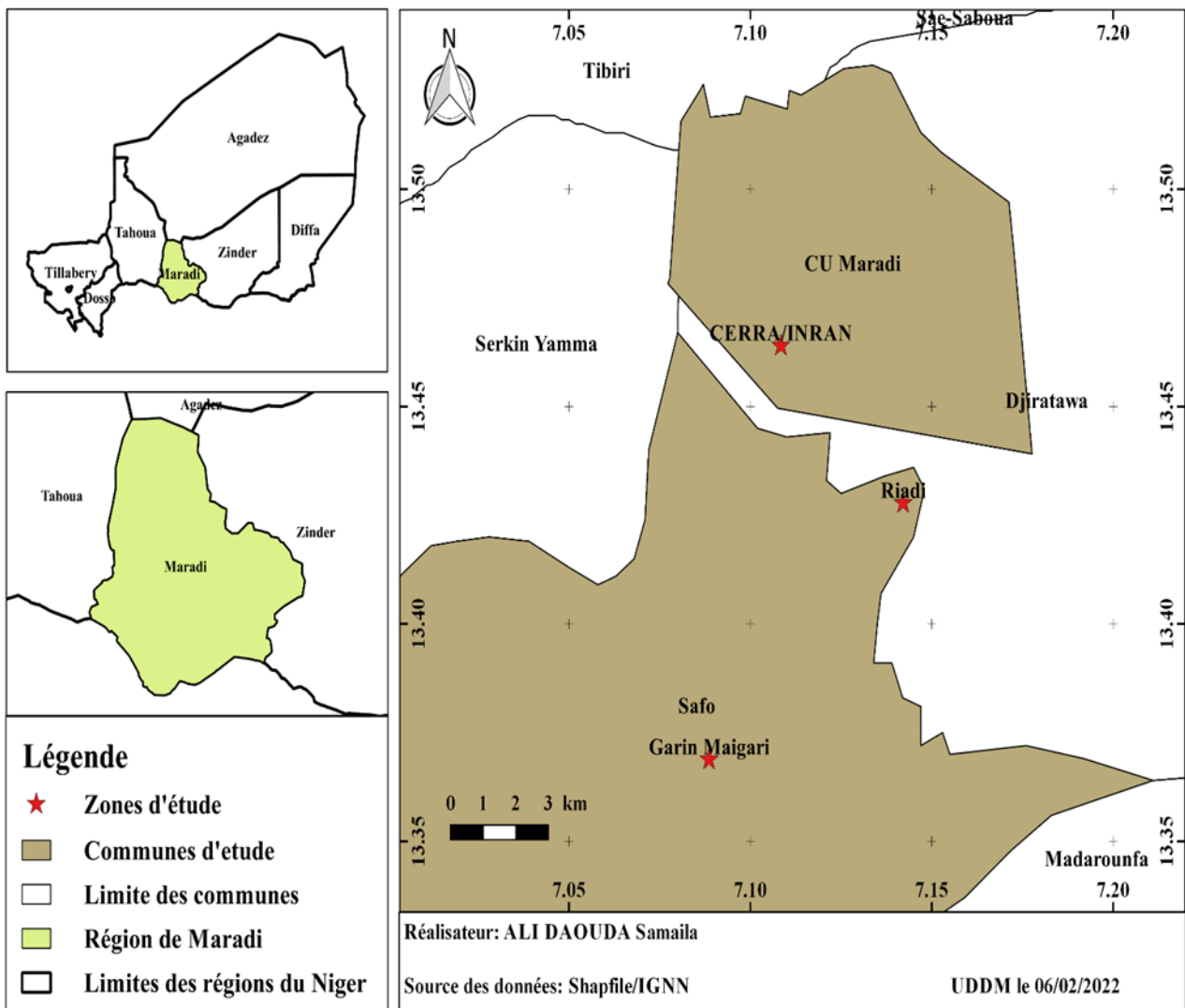


Figure 1: Representative map of the different experimental sites

### 3. Experimental design

The same experimental design was used on all sites. It is a randomized incomplete block design with three repetitions. Each repetition consists of 6 blocks of 15 m<sup>2</sup> which is made up of 5 varieties. The spacings between the lines are 0.80 m and 0.30 m between hills.

### 4. Data analysis

The data obtained were analyzed by the GENSTAT (software version 15). This software allowed to carry out the analysis of variance with the student test for the comparison of the means of the different varieties. A correlation matrix using the same software for all of the measured parameters was produced. It also allowed through GGE Biplot analysis methods, to evaluate the genotype x environment (GxE) interaction and identify the most efficient genotypes on the one hand and the most stable on the other.

**Data collected were:** VAR: Varieties; NPL: Number of hills Raised; 50% flo: 50% Flowering; HTR: Plant Height; MAT: Maturity physiology; NPR: Number of Panicles Harvested; PP: Panicle weight; RDT G: Seed yield; RTD T: Stem Yield; IR: Harvest Index; IB: Threshing Index, T1S: Local Witness 1 Station; T2S: Local indicator 2 Station; JAS = Days After Semi

## III. RESULTS

### 1. CERRA Maradi Station

At this site, for the 50% Flo variable, the TLS1 (MM) variety flowered very early with an average of 64.33 DAS and the V49 variety the latest with an average of 103.6 DAS. For the maturity variable (MAT), variety V49 recorded the highest average (118.7 DAS) and variety TLS1 recorded the lowest average (79.3 DAS).

In other words, at this site, the TLS1 variety (MM) was found to be the earliest with (79.3 DAS) while V49 was the latest with (118.7 DAS) (table 1). On the other hand, the averages of the other varieties are comparable. For this site no significant difference was observed for plant height (HTR). Regarding grain yields, the MDK variety is the most efficient with a yield of 1333.3 kg/ha while the V49 has the lowest yield (111.1 kg/ha). Indeed, for the average stem yield (RDT T) the MDK variety has the highest average with 5444 kg/ha and that of the V29 varieties the lowest (2831 kg/ha). For the variable number of panicles harvested (NPR), variety SSD 35 recorded the highest average (21 panicles) and V49 the lowest (2.67 panicles). Regarding the panicle weight (PP) variable, the MDK variety recorded the highest average (1.05 kg) and V49 the lowest (0.08 kg).

**Table 1: Performance of Sorghum varieties at CERRA Maradi**

VAR	50% flo	HTR	MAT	NPR	PP	RDT G	POI100 GR	RDT T	IR	IB
V49	103,6a	130a	118,7a	2,67c	0,08b	111,1a	4b	5667ab	2,70bc	0,44ab
L 28	102,67a	133,3a	117,7a	11,33abc	0,20b	777,8a	4ab	6256ab	2,87bc	1,91ab
V73	102,00a	123,3a	117a	6abc	0,10b	166,7a	4b	5067ab	2,52bc	0,50ab
V39	101,33a	101,7a	116,3a	16,67abc	0,21b	388,8a	4b	6917ab	3,07bc	0,55ab
V68	101,00a	111,7a	116a	4,33bc	0,16b	333,3a	4,33ab	6572ab	2,97bc	0,61ab
V33	99,67a	156,7a	114,7a	10,67abc	0,15b	250a	3,33b	12217bc	4,66ab	0,50ab
MDK	98,67a	116,7a	113,7a	20a	1,05a	1333,3b	6a	15444c	5,63a	0,39ab
V17	98,33a	93,3a	113,3a	15,67abc	0,26b	444,4a	4b	6556ab	2,96bc	0,50ab
V61	97,33a	146,7a	112,3a	7abc	0,10b	194,4a	2,66b	5222ab	2,56bc	0,55ab
V8	97,00a	128,3a	112a	8,33abc	0,15b	227,8a	4,66ab	8972ab	3,69bc	0,55ab

V45	96,67a	130a	111,7a	12,67abc	0,33b	388,9a	4,66ab	7989ab	3,39bc	0,36ab
V19	96,33a	143,5a	111,3a	10abc	0,13b	250a	4b	8683ab	3,60bc	0,55ab
V40	96,33a	91,7a	111,3a	6,67abc	0,10b	166,7a	3,66b	5111ab	2,53bc	0,50ab
V47	95,67a	120a	110,7a	11abc	0,16b	227,8a	4,33ab	2556abc	4,16abc	0,50ab
V1	95,33a	110a	110,3a	10,33abc	0,25b	361,1a	4,66b	6017ab	2,80bc	0,46ab
V27	94,67a	100a	109,7a	10abc	0,16b	333,3a	4,66b	7861ab	3,36bc	0,61ab
V13	94,00a	130a	109a	10abc	0,15b	227,8a	3,66b	4000a	2,20c	0,55ab
V3	93,67a	110a	108,7a	14,33abc	0,18b	333,3a	4,66ab	8967ab	3,69bc	0,55ab
V48	93,67a	158,3a	108,7a	7abc	0,11b	222,2a	4b	6111ab	2,83bc	0,66a
V29	93,33a	86,7a	108,3a	7,33abc	0,10b	166,7a	4,33ab	2831a	1,85c	0,50ab
V70	91,00ab	106,7a	106ab	8,67abc	0,18b	227,8a	4,33ab	5333ab	2,60bc	0,46ab
V37	87,00ab	100a	102ab	18,67ab	0,26b	388,9a	3,66b	7556ab	3,26bc	0,48ab
V16	85,67ab	126,7a	100,7ab	9abc	0,16b	333,3a	4b	4494ab	2,35bc	0,61ab
V5	85,67ab	115a	100,7ab	7,67abc	0,11b	694,4a	2,66b	4344ab	2,30bc	2a
V50	84,33ab	141,7a	99,3ab	12abc	0,15b	227,8a	3,66b	5000ab	2,50bc	0,55ab
L153-5	83,33ab	115a	98,3ab	8,67abc	0,10b	166,7a	3,33b	2972a	1,89c	0,50ab
V58	83,00ab	145a	98ab	17,67abc	0,26b	444,4a	3,66b	9222ab	3,76bc	0,50ab
TL2S	74,67bc	135a	89,7bc	16abc	0,21b	333,3a	3,66b	4778ab	2,43bc	0,51ab
SSD35	65,00c	136,7a	80c	21a	0,25b	388,9a	3b	6350ab	2,90bc	0,51ab
TL1S	64,33c	135a	79,3c	12abc	0,15b	227,8a	3,66b	3111a	1,93c	0,55ab

Means followed by the same letter in the columns are not significantly different according to the LSD test at 5% probability

## 2. RIADI

At the village of Riadi, dates of 50% flowering (50% FLO) of varieties are between 62.67 DAS and 98.33 DAS (table 2). The variety SSD 35 with 62 DAS is the first variety to reach this stage while the MDK and TL1R (MDW) varieties with 98.33 DAS represent the last. Physiological maturity (MAT) of varieties is reached between 77.67 and 133.33 DAS (table 2). This physiological maturity differs significantly depending on the variety. Indeed, SSD 35 with 77.67 DAS is the earliest variety while MDK with 133.33 DAS is the latest.

For the height parameter (HTR), the TL1R (MDW) variety recorded the highest average (265 cm), the varieties V73, V68, MDK, V8, V58, V5, SSD 35 are part of the same group. The V40 recorded the lowest average (110 cm). There is no significant difference between the 30 lines with regard to the NPR and PP parameters. For grain yield (RDT G), it differs significantly. The varieties V27 and TL1R have the highest yields with 1388.7 kg/ha and 1222.2 kg/ha respectively while V33 and V37 gave the lowest yields (333.3 kg/ha). In terms of stem yield (RDT T) (Table 2), the analysis revealed a significant difference between the varieties. The MDK and TL2R varieties have very high yields with 8111kg/ha and 7889kg/ha respectively while the L153-5 gave the lowest stem yield (2222kg/ha).

**Table 2: Performance of sorghum varieties at RIADI**

VAR	50% flo	HTR	MAT	NPR	PP	RDT G	POI100 GR	RDT T	IR	IB
V49	87,33bc	156,7defghi	102,33bc	17,77	0,45	500abcd	4bcd	3111a	1,93c	0,35abc
L 28	85,67bcd	160defghi	100,67bcd	16,33	0,40	444,4abc	4,66b	5000abc	2,50abc	0,32bc
V73	82,67bcdef	170cdefgh	97,67bcdef	16	0,73	833,3abcd	4bcd	3778ab	2,13bc	0,36abc
V39	85bcd	140dfghi	100bcd	20	0,66	777,8abcd	4,33bc	3667ab	2,10bc	0,35bc
V68	81,67bcdefg	186,7cdef	96,67bcdefg	14,67	0,38	444,4ab	4,33bc	4000abc	2,20abc	0,35abc
V33	84,67bcd	158,3defghi	99,67bcd	15,33	0,35	333,3a	4,33bc	5222abc	2,56abc	0,31bc
MDK	98,33a	240ab	113,33a	17,33	0,61	722,2abcd	4bcdef	8111c	3,43a	0,32bc
V17	80,33bcdefg	121,7ghi	95,33bcdefg	21,67	0,85	1000abcd	3,33cdef	3222a	1,96c	0,39bc
V61	74,33defgh	131,7fghi	89,33defg	18,33	0,8	833,3abcd	4bcd	5333abc	2,60abc	0,31bc
V8	77,67bcdefg	188,3cde	92,67bcdefg	20,33	0,50	500abcd	4,66b	4667abc	2,48abc	0,27bc
V45	87,67b	130fghi	102,67b	14,33	0,46	444,4ab	4bcd	2889a	1,86c	0,31bc
V19	80,33bcdefg	125ghi	95,33bcdefg	19,67	0,50	555,6abcd	4bcd	3222a	1,96c	0,33bc
V40	75defg	110i	90,02defg	12,67	0,50	500abcd	3,33cdef	4111abc	2,23abc	0,28bc
V47	82,67bcde	213,3bc	97,67bcde	17,33	0,46	666,7abcd	4,66b	4444abc	2,33abc	0,42bc
V1	85,33bcd	143,3defghi	100,33bcd	17,33	0,31	388,9a	4bcd	3222a	1,96c	0,35abc
V27	80,33bcdefg	153,3defgh	95,33bcdefg	18,33	0,68	1388,9bd	4,33bc	5556abc	2,66abc	0,60a
V13	74,33defgh	113,3hi	89,33defg	20,67	0,78	444,4ab	3dg	4222abc	2,26abc	0,20c
V3	84,67bcd	161,7defghi	99,67bcd	18	0,63	666,7abcd	4bcd	4556abc	2,36abc	0,32bc
V48	87,33bc	180cdefg	101,33bc	17,77	0,56	555,6abcd	433bc	3889abc	2,16abc	0,29bc
V29	71,67efghij	131,7fghi	86,67efghij	12	0,31	388,9a	4bcd	5000abc	2,50abc	0,36abc
V70	76,33bcdefg	120hi	91,33bcdefg	16,67	0,50	444,4ab	4bcd	4778abc	2,43abc	0,28bc
V37	70,67efghijk	128,3ghi	85,67efghijk	17	0,50	333,3a	3,66bcdefg	4556abc	2,36abc	0,20c
V16	74defgh	126,7ghi	80,2efg	20,33	0,63	777,8abcd	4bcd	5222abc	2,56abc	0,35abc
V5	75,67cdefg	165defghi	90,67cdefg	13	0,68	444,4ab	3,33cdef	4778abc	2,43abc	0,23c
V50	73,67defghi	140dfghi	88,67defghi	17,67	0,56	722,2abcd	4bcd	4222abc	2,26abc	0,39abc
L153-5	64,33hijkl	143,3defghi	79,33hijkl	18,33	0,55	500abcd	4bcde	2222a	1,66c	0,37bc
V58	75,33defg	131,7efghi	90,33defg	16,67	0,66	833,3abd	4bcd	5667abc	2,70abc	0,35abc
TL2R	71efghij	260a	86efghij	16	0,58	1055,6abcd	3dge	7889bc	3,36abc	0,55ab
SSD35	62,67jl	193,3cd	77,67jl	24,67	0,71	944,4abcd	3,66bcdefg	3333a	2c	0,39abc
TL1R	98,33a	265a	113,33a	24,33	0,95	1222,2abc	6a	5556abc	2,96abc	0,40bc

Means followed by the same letter in the columns are not significantly different according to the LSD test at 5% probability.

### 3. GARIN MAIGARI

At the Garin Maigari level, there is no significant difference regarding the plant height parameters (HTR), stem yield (RDT T) and panicle weight (PP). The other parameters differ statistically depending on the varieties. For the 50% FLO and MAT parameters, the TL1G variety (El Bazanga) is the earliest with 84 DAS and the MDK is the latest with 107 DAS. For the numbers of panicles harvested (NPR), the SSD 35 variety gave the highest number with 26 panicles, the TLG2 and V70 varieties gave the same number of panicles (23 panicles) while the V5 (8.33 panicles) and V40 (9 panicles) gave the lowest numbers of panicles. The varieties MDK and TLG2 (MDW) gave the best grain yield (1111 kg/ha) while L153-5 was the least efficient with a yield of 222.2 kg/ha (Table 3). The average weight of 100 grains of the TLG2 and MDW varieties (66g) are the highest and those of L 153-5, V49, V1 (3.33g) the lowest.



**Table 3: Performance of sorghum varieties at Garin Maigari**

VAR	50% flo	HTR	MAT	NPR	PP	RDT G	POI100 GR	RDT T	IR	IB
V49	104,33ab	140a	119,3ab	12ab	0,23b	277,8a	3,33b	6222a	2,86a	0,33ab
L 28	101,33ab	113,3a	116,3	11,67ab	0,21b	500a	4b	4111a	2,23a	1,91a
V73	100,67abcd	128,3a	115,7abcd	8,67b	0,16b	277,8a	4b	10556a	4,16a	0,50b
V39	96,67abcd	110a	111,7abcd	16,33ab	0,35b	444,4a	4b	7333a	3,20	0,36b
V68	96abcd	143,3a	111abcd	16,67ab	0,35b	388,9a	4b	10556a	3,16a	0,33ab
V33	99,33abcd	121,7a	114,3abcd	17ab	0,28b	277,8a	4,33ab	6667a	3a	0,36b
MDK	107a	178,3a	122a	19,67ab	1,20a	1111,1b	4,66ab	5889a	4,5a	0,28b
V17	98abcd	145a	113abcd	18,67ab	0,56b	500a	4,33b	11667a	2,76a	0,26b
V61	88,33bcd	133,3a	103,3bcd	16ab	0,30b	444,4a	4b	6000a	2,80a	0,46b
V8	94abcd	153,3a	109abcd	20,33ab	0,46b	555,6a	4,66ab	9444a	3,83a	0,36b
V45	101abc	110a	116abc	16ab	0,48b	500a	4,66ab	6111a	2,83a	0,35ab
V19	88,33bcd	170a	103,3bcd	17ab	0,41b	611,1a	4b	6111a	2,83a	0,45b
V40	98,67abcd	131,7a	113,7abcd	9b	0,18b	333,3a	3,66b	3000a	1,9a	0,66b
V47	97,33abcd	163,3a	112,3abcd	13ab	0,23b	333,3a	4b	10556a	4,16a	0,44b
V1	103,67abc	108,3a	118,7abc	10b	0,2b	277,8a	3,33b	5444a	2,63a	0,57b
V27	94abcd	118,3a	109abcd	14ab	0,28b	444,4a	3,66b	3333a	2a	0,47b
V13	93,33abc	116,7a	108,3abcd	11,67ab	0,18b	277,8a	3,66b	5556a	2,66a	0,44b
V3	97,33abc	136,7a	123,3abcd	16ab	0,41b	555,6a	4b	8889a	3,66	0,38b
V48	98abcd	160a	113abcd	15,33ab	0,25b	277,8a	4b	5889a	2,76a	0,52b
V29	87,67bcd	128,3a	102,7bcd	12,67ab	0,2b	277,8a	4b	7667a	3,30a	0,44b
V70	90,67abcd	125a	105,7abcd	23ab	0,55b	722,2a	4b	8111a	3,34a	0,38b
V37	89bcd	11,7a	104bcd	21ab	0,33b	333,3a	4b	4444a	2,33a	0,33ab
V16	90bcd	128,3a	105bcd	19ab	0,41b	555,6a	4,33ab	6333a	2,90a	0,41b
V5	94,67abcd	165a	109,7abcd	8,33b	0,23b	277,8a	4b	4111a	2,23a	0,36ab
V50	87,67bcd	116,7a	102,7bcd	18,33ab	0,21b	277,8a	4,33ab	4889a	2,46a	0,38b
L153-5	87cd	116,7a	102cd	16,67	0,13b	222,2a	3,33b	3778a	2,13a	0,50b
V58	89,67bcd	121,7a	104,7bcd	13,33ab	0,25b	500a	4b	3111a	1,93a	0,59b
TL2G	101,67abc	183,3a	116,7abc	23ab	1,10a	1111,1b	5,66a	11111a	4,3a	0,30b
SSD35	92abcd	155a	107abcd	26a	0,31b	500a	4b	7778a	3,33a	0,48b
TL1G	84d	156,7a	99d	17,67ab	0,30b	388,9a	5ab	7889a	3,36a	0,39b

Means followed by the same letter in the columns are not significantly different according to the LSD test at 5% probability.

**4. Combined analysis results from the 3 sites**

The combined analysis of variance (Table 4) shows that there is a significant interaction between sites and varieties for six (6) parameters, namely: flowering 50% (50% Flo), plant height (HTR), physiological maturity (MAT), panicle weight (PP), grain yield (RDT GR) and 100 grain weight (POI 100 GR).

**Table 4: variances of the different parameters monitored**

Sources of variance	50% flo	HTR	MAT	NPR	PP	RDT GR	POI100 GR	RDT T	IR	IB
Variety (V)	10,54*	4,85*	10,54*	4,19*	5,28*	3,88*	4,47*	2,96*	2,93*	1,37ns
Sites (S)	167,60*	35,27*	167,60*	47,05*	92,75*	30,40*	1,15ns	17,12*	17,13*	6,62*
V x S	3,46*	1,83*	3,46*	1,27ns	2,28*	1,80*	2,91*	1,39ns	1,39ns	0,68ns

'ns' not significant, '\*' significant

**5. Correlation between the studied parameters**

At this level the correlation matrix between the parameters which allows us to see the link between the pairs of variables has been highlighted (Table 5). Indeed, a strong significant correlation is observed between the RDT G variable and the HTR with (r=0.29). The correlation between RDT G and RDT T is highly significant and positive with (r=0.15). The correlation between the RDT T and the NPR is very highly significant with a coefficient (r=0.21). The correlation between RDT G and MAT is highly significant and negative with (r=-0.15). The correlation between the RDT G and the NPR, as well as that

observed between the RDT G and the PP, are also very highly significant with respective coefficients (r=0.48) and (r=0.75).

**Table 5: Correlation matrix between variables**

	50%Flo	HTR	IB	IR	MAT	NPR	POI 100G	PP	RDT G	RD T T
50%Flo	1									
HTR	-0,131*	1								
IB	0,143*	-0.095ns	1							
IR	0.227***	0.089ns	-0.076ns	1						
MAT	1.000***	-0.131*	0.1430*	0.227***	1					
NPR	-0.308***	0.218***	-0.274***	0.216***	-0.308***	1				
POI 100G	0.193***	0.099ns	-0.099ns	0.269***	0.193***	0.232***	1			
PP	-0.234***	0.315***	-0.245***	0.160**	-0.234***	0.617***	0.323***	1		
RDT G	-0.158**	0.299***	0.243***	0.158**	-0.158**	0.481***	0.286***	0.757***	1	
RDT T	0.227***	0.089ns	-0.076ns	1.000***	0.227***	0.216***	0.269***	0.160**	0.158**	1

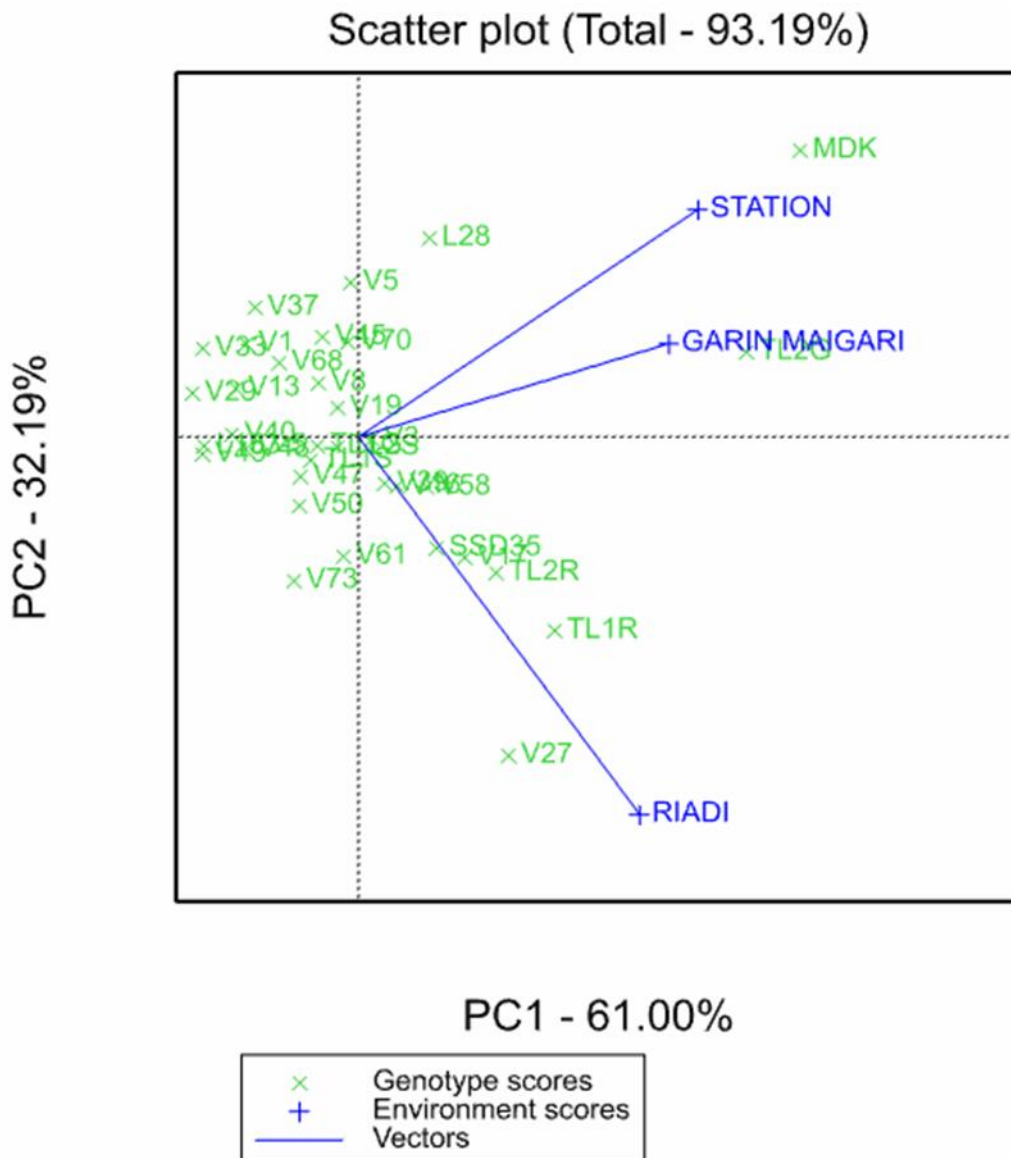
‘ns’ not significant ‘\*\*\*’ significant at 0.1%; ‘\*\*’ significant at 1%; ‘\*’ significant at the 5% level.

## 6. GGE Biplot Analysis

### 6.1. Environment Vector Plot

The results showed a distribution of environments and genotypes following the two main components PC1 and PC2. The environments are represented by vectors in blue color and the genotypes by their names in green color. The angle formed by the environment vectors provides an estimate of their similarity. Analysis of the figure reveals that for grain yield, the vectors of the CERRA Maradi STATION and RIADI environments form an obtuse angle. These two environments are therefore totally different, on the other hand the CERRA Maradi and GARIN MAIGARI environments form an acute angle (less than 90°). They are similar and form a mega environment

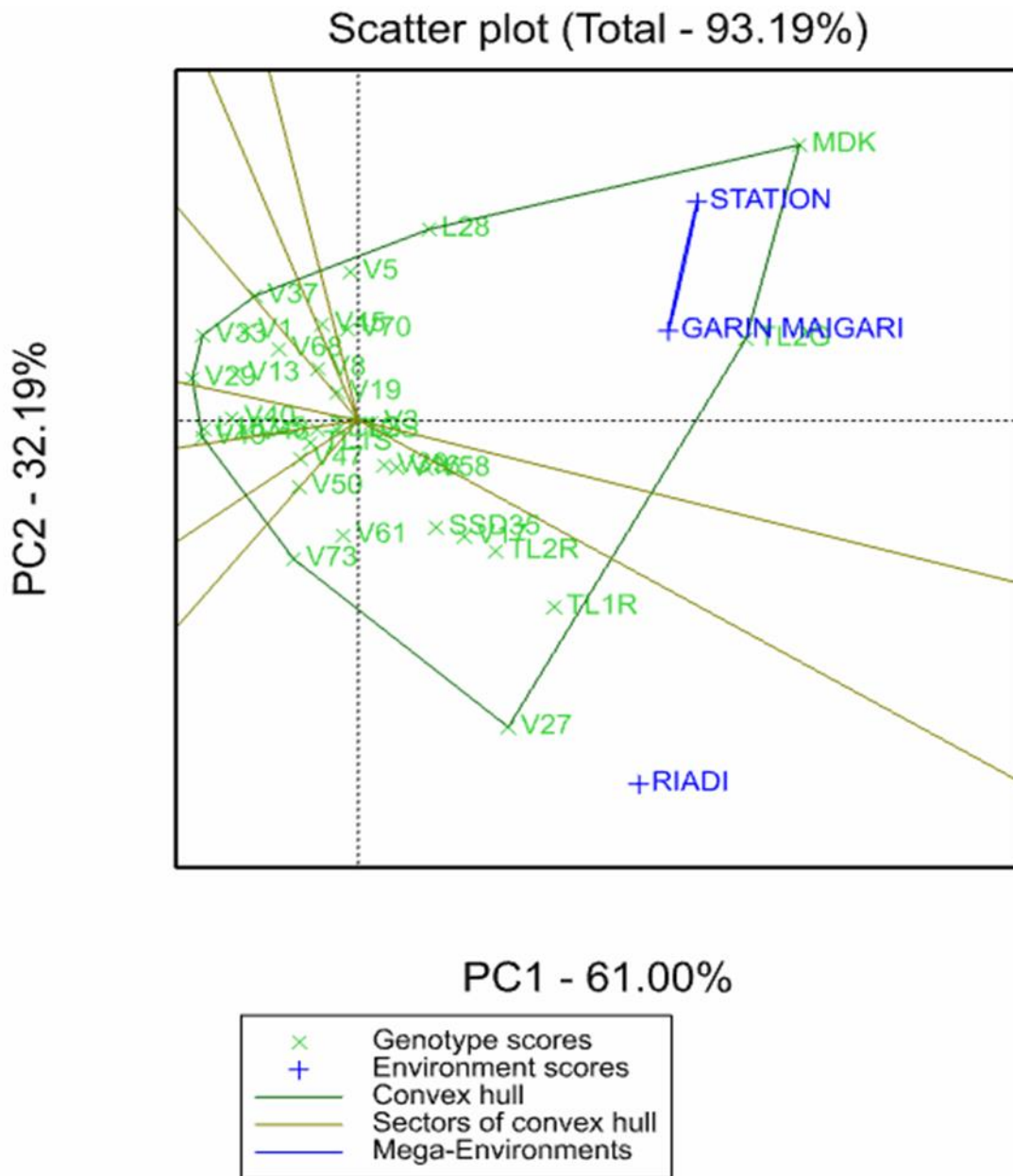




**Figure 2: Distribution of genotypes and environments for Grain Yield**

### 6.2. Expression of genotypes depending on environments

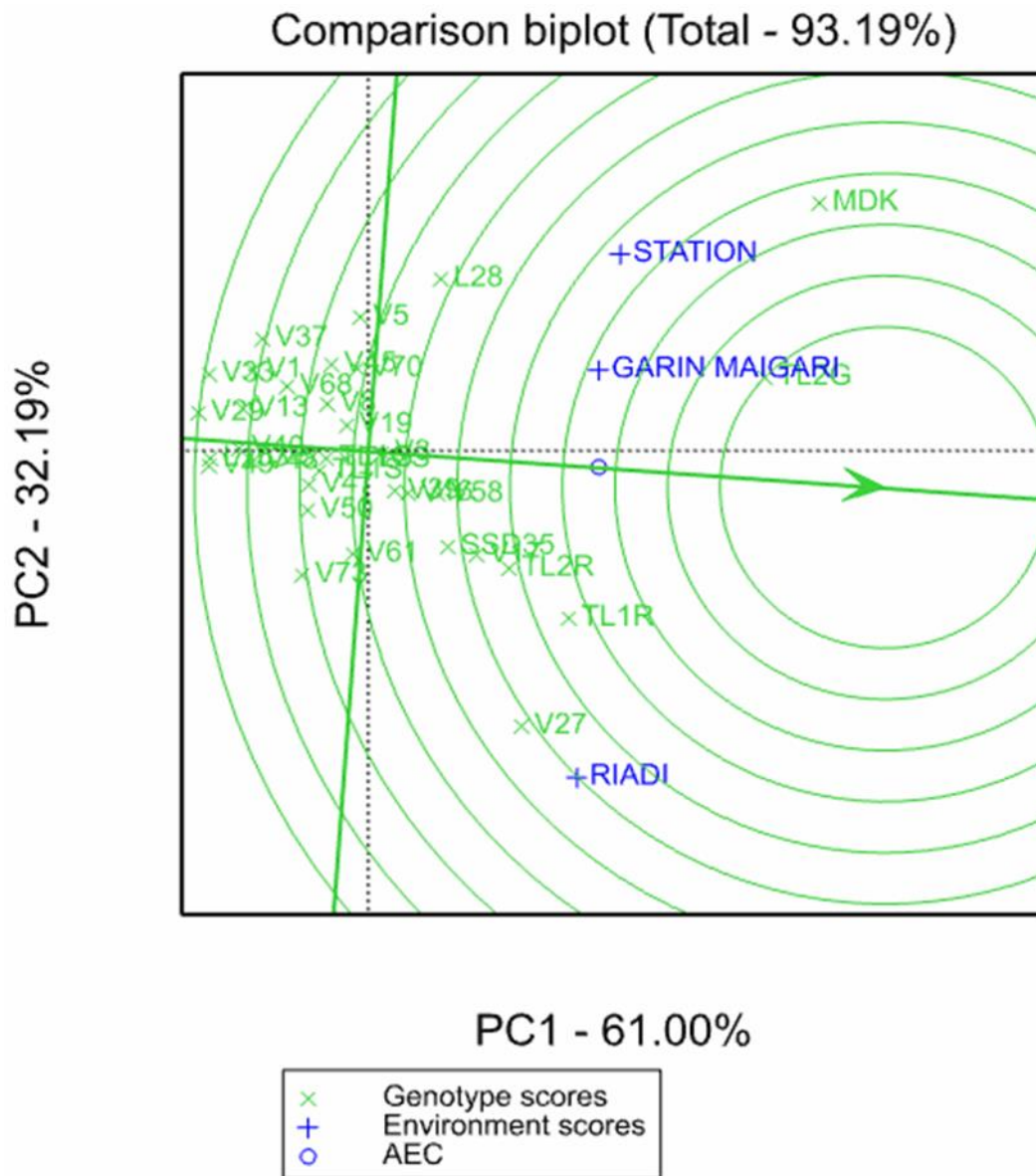
The projection of the genotypes onto the plane formed by the two principal components PC1 and PC2 gives a polygon whose vertices are constructed by the elite genotypes. The polygon is subdivided into several compartments by straight lines perpendicular to the polygon forming the mega-environments. Genotypes found in an area including an environment are best suited to that environment. These are the MDK and TL2G genotypes which are the most efficient in the mega environment constituted by STATION and GARIN MAIGARI, the V27, TL1R and V73 genotype in the RIADI environment.



**Figure 3: Distribution of genotypes, environments and mega-environment for Grain Yield (RDT G)**

### 6.3. Comparison of genotypes

For the comparison of genotypes in terms of grain yield (RDT G) according to this analysis, the ideal genotype (which has the highest and stable average) is the one which is closer to the circle. Thus, the further we move away from the circle, the less efficient the genotype is. This is the TL2G genotype which has the highest average in terms of grain yield.



**Figure 4: Comparison of genotypes**

#### 6.4. Comparison of environments

This analysis makes it possible to measure the representativeness of the environments and to define a benchmark environment and to use it as a reference for comparison, this “ideal” environment, or the average environment (indicated by a circle with an arrow in it representing the axis AEC). This environment is the most discriminating of the genotypes and the most representative compared to other environments. in the present study it is GARIN MAIGARI which is the most desirable for evaluating the performance of the genotypes.

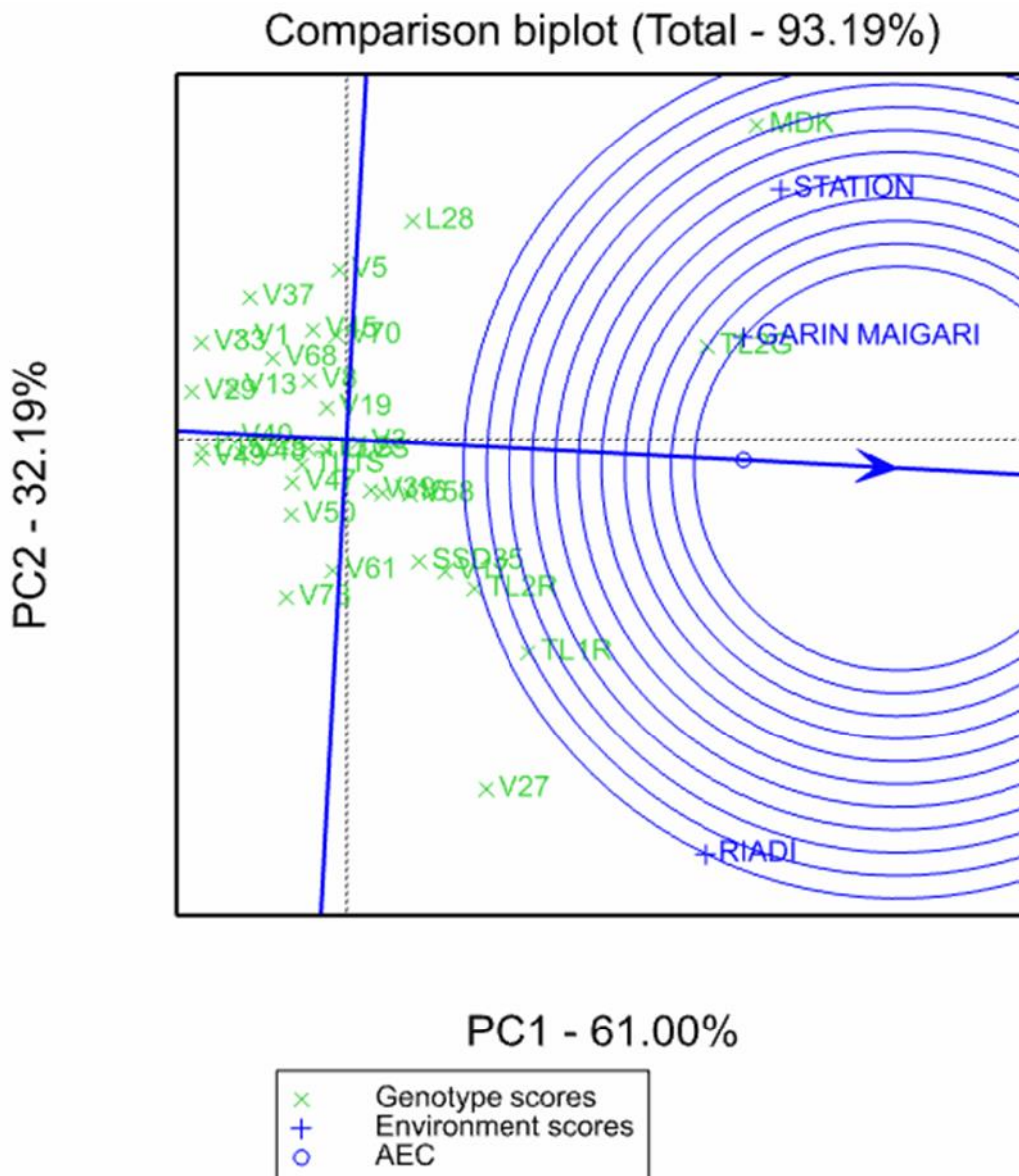
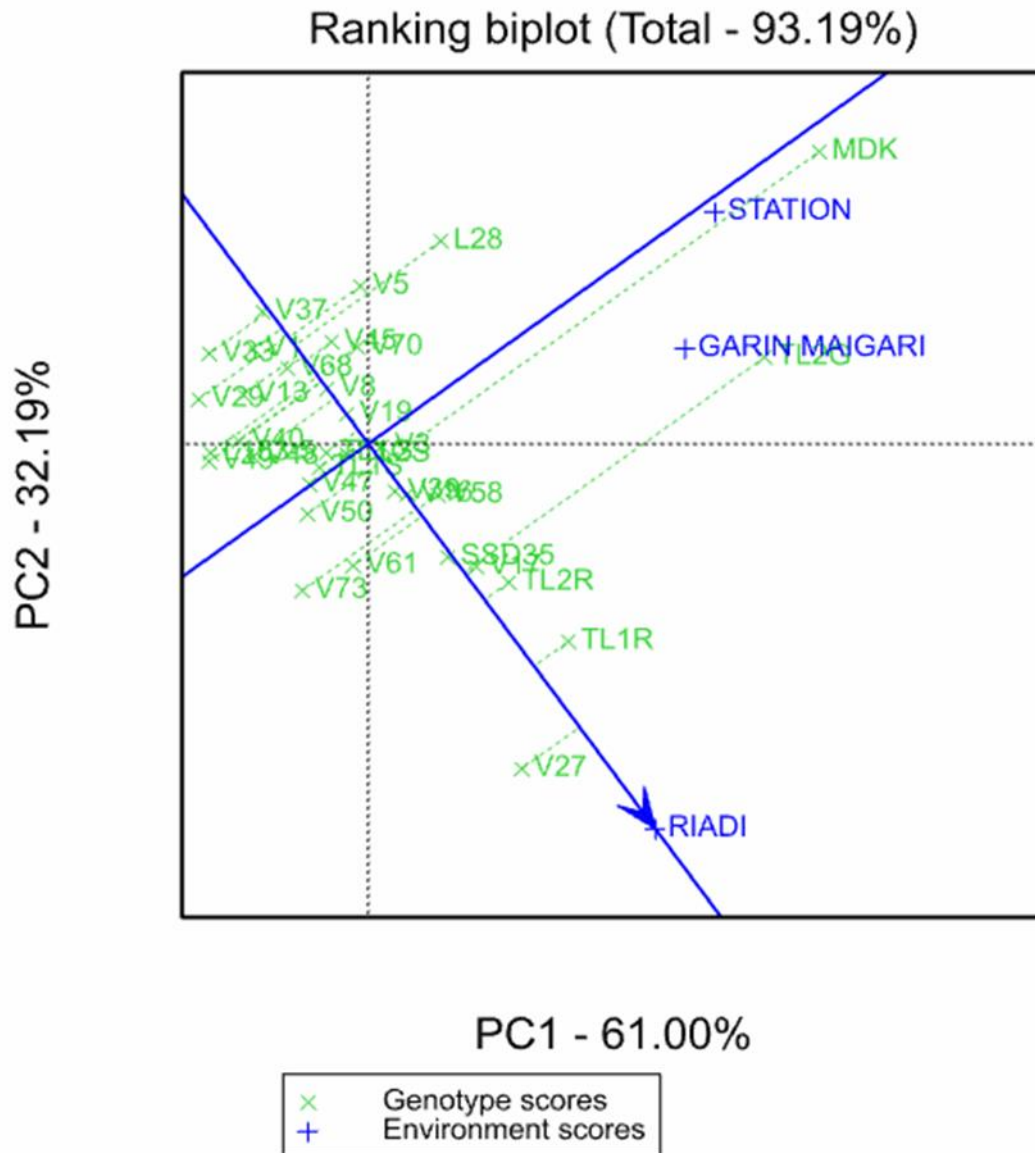


Figure 5: classification of environments according to representativeness and discriminating capacity.

### 6.5. Classification of genotypes according to environment

This analysis makes it possible to classify varieties according to their performance in an environment. The distance between axis and genotypes determines genotype performance in this environment. However, the smaller the distance between the axis and the genotype, the more efficient the genotype is in this site. For the RIADI site, the most efficient genotypes are among others V27, TL1R, TL2R, SSD 35, V61.



**Figure 6: Classification of genotypes according to environment**

#### IV DISCUSSION

The results of the present study indicate a very large variability for the measured parameters. This variability is both genetic and environmental. The descriptive analysis confirms this variability, where we see that the variables differ between genotypes and between sites. Indeed, the analysis of variance for each site showed us a significant difference between the varieties for all parameters. The significance that exists between the majority of parameters for the different localities allows us to deduce that the different sites are varied.

On the other hand, the significant difference for all the characteristics studied with the exception of the number of panicles harvested, the stem yield, the harvest index and the threshing index for the interaction between environment and varieties reflects a particular influence for characters as mentioned by Gaufichon et al., 2010. He also indicated that the expression of genes can be modified depending on environmental conditions confirmed by the studies carried out by Garba et al., (2015) thus attesting that the varieties are generally developed based on traits that allow them to adapt to their environment. These



traits are closely linked to the ecological conditions in the production areas. In other words, the variation observed for the traits measured is partly due to environmental factors coupled with genotypic differences.

Regarding correlation, it is very useful for selection work according to Siene et al. 2010. For our study, the analysis of the correlations between the measured parameters shows that many variables presented an interesting correlation from the morphological and agronomic point of view. However, we find between the grain yield variable (RDT G) and plant height (HTP), a highly significant and positive correlation. The latter explains that the higher the stems of the plant, the higher the grain yield. Likewise, a highly significant and positive correlation is observed between grain yield (RDT G) and stem yield (RDT T). This shows that the more stem biomass the plant produces, the more grain yield it has. The link can also explain the plant's ability to transfer assimilates to grain production. This positive correlation between the RDT G and the RDT T makes it possible to select genotypes which have a good ability to transfer assimilates for grain production. We also observe a positive correlation between the variable Height of plants (HTP) and 50% Flowering (50%FLO) which allows us to understand that precocity depends on the height of the plant in this study.

Late varieties have a higher height since the plant will take a long time to complete its cycle and therefore will have time for good vegetative development. This result is consistent with that of Illiassou, 2017 where he also found a significant and positive correlation between HTP and FLO.

There is a non-significant correlation between stem yield (RDT T) and plant height (HTP). This allows us to understand that for the present study, in sorghum, stem yield is not only linked to the height of the plants but a combination of several parameters such as stem diameter and number of leaves. This result contradicts that of Illiassou, 2017 where he found a significant and positive correlation between RDT T and HTP.

In other words, the results of the GGE Biplot analysis rank genotypes by their average yield performance and stability in a number of environments. An ideal genotype is one that is efficient and stable. In our study, the most efficient and stable genotypes are therefore V68, V3, V58. This result is in the same direction as that of Naroui et al. (2013) on wheat who found genotypes (14, 20, 21, and 22) more stable with appropriate yield.

This result shows that the effect of the environment plays a very important role in the expression of yield of certain varieties, which is why it is rare to find genotypes expressing their potential in several different environments. This is the case of a study on 10 sesame genotypes which highlighted a single GK01 genotype as the best cultivar for the 5 test environments (Boureima, 2017).

The genotypes SSD 35, V61, L28, V27, MDK have average high yields but they are not stable and the genotypes V 13, V 47, V 43, V19, V29, V8 are stable but very low yields (lower to the average).

These results corroborate those of Ménad et al. (2010) which stipulate that the stability of yields is independent of their values, and that genotypes with high yield are generally relatively unstable while it is generally the opposite for genotypes with medium or low yield. They also confirm the assertion of Yan and Hunt (2002) who indicate that overall stability is not necessarily a positive factor and that it is only desirable when it associates a high average return. (Ndiaye et al., n.d.).



## V. CONCLUSION

Sorghum is a critically important food crop worldwide. This crop is exposed to different factors that influence its production such as drought, diseases, pest attacks, lack or inaccessibility of improved varieties.

Varietal innovations have always been a key element of agricultural production. Evaluating the behavior of new varieties in the face of a diversity of environments is therefore an important issue for stakeholders, from those who select these new varieties to those who cultivate them or use the products.

Our study made it possible to evaluate agronomic performances depending on the sites. And the GGE BILOT analysis showed that grain yields were strongly influenced by genotypes, environments and genotype-environment interaction. Environments elicited different responses from genotypes and most of these genotypes showed environmental specificity.

This work carried out during the 2021 rainy season in the villages of RIADI, GARIN MAIGARI and at the CERRA-Maradi STATION noted very appreciable abilities of certain genotypes with regard to agro morphological variables and yields. The GGE Biplot analysis made it possible to identify mega environments. However, two mega-environments are determined for the grain yield variable. GARIN MAIGARI and STATION are similar and different from the RIADI environment. The genotypes most suitable for the mega-environment formed by RIADI and GARIN MAIGARI are MDK, TL2G, L28 and V5 while V27 and V73 are the best genotypes in the RIADI environment. This analysis also showed that GARIN MAIGARI constitutes the ideal environment (benchmark) and can therefore be used for genotypes with broad adaptation.

In addition, this analysis made it possible to identify certain genotypes such as SSD 35, V61, L28, V27, MDK which have high but unstable average yields and V 13, V 47, V 43, V19, V29, V8 which are stable but producing very low yields (below average).

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